



ROWERSA CORPORATION CALLEN, Walter MATHUR, Eric

<120> ENZYMES HAVING HIGH TEMPERATURE POLYMERASE ACTIVITY AND METHODS OF USE THEREOF

DIVER1350-2															
<140> US 09/656,309 <141> 2000-09-06															
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<170> PatentIn version 3.0															
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ggt aaa gag cct cag gta atc ata tgg ggt att gct gag aac ggc gag Gly Lys Glu Pro Gln Val Ile Ile Trp Gly Ile Ala Glu Asn Gly Glu 20 25 30	96														
agg gta gtc ctc att gac agg tct ttt cgc cca tac ttc tat gcg ctg Arg Val Val Leu Ile Asp Arg Ser Phe Arg Pro Tyr Phe Tyr Ala Leu 35 40 45	144														
ctt gca ccg ggc gcc gat cct aag cag gta gca caa cgt att cgt gca Leu Ala Pro Gly Ala Asp Pro Lys Gln Val Ala Gln Arg Ile Arg Ala 50 55 60	192														
ttg agt agg cca aag agc ccg att ata ggt gta gag gat gac aag agg Leu Ser Arg Pro Lys Ser Pro Ile Ile Gly Val Glu Asp Asp Lys Arg 65 70 75 80	240														
aag tac ttc ggg agg cct cgt agg gtc tta cgt att cgc acc gtg cta Lys Tyr Phe Gly Arg Pro Arg Arg Val Leu Arg Ile Arg Thr Val Leu 85 90 95	288														
ccc gag gct gtt agg gag tat cgc gaa ctc gta aag aac gtt gat ggt Pro Glu Ala Val Arg Glu Tyr Arg Glu Leu Val Lys Asn Val Asp Gly 100 105 110	336														
gtt gag gat gtt cta gag gcg gat ata cgc ttc gct atg cgc tat ctc	384														

13

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Val	Glu	Asp 115	Val	Leu	Glu	Ala	Asp 120	Ile	Arg	Phe	Ala	Met 125	Arg	Tyr	Leu	
										tac Tyr						432
				_	_			-	_	gac Asp 155	_	_		_	_	480
										gct Ala						528
										att Ile						576
										gta Val						624
										att Ile						672
										tac Tyr 235						720
										cat His						768
										aag Lys						816
					Pro		Thr	Ser		cat His		His	Val			864
										tat Tyr						912
gag Glu 305	atc Ile	aag Lys	ata Ile	aag Lys	agt Ser 310	ctc Leu	gag Glu	gag Glu	gtc Val	gca Ala 315	gag Glu	tat Tyr	cta Leu	ggc Gly	gtg Val 320	960
										tgg Trp						1008
										tta Leu						1056

									aag Lys				1104
									cta Leu 380				1152
									ctg Leu				1200
	_	_			_		_	_	gag Glu	_	_		1248
									aga Arg				1296
									cca Pro				1344
_			 	_	_				cct Pro 460				1392
		_	 _	 	_	_		_	aag Lys			-	1440
	-	_							gag Glu				1488
									tat Tyr				1536
									gcg Ala				1584
									ggc Gly 540				1632
									ggt Gly				1680
									ctc Leu				1728

												gag Glu				1776		
aat Asn	ttc Phe	atc Ile 595	aaa Lys	att Ile	ata Ile	aag Lys	gag Glu 600	gag Glu	ctg Leu	gly aaa	ttc Phe	gaa Glu 605	atc Ile	aag Lys	cta Leu	1824		
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												ggt Gly				1920		
gta Val	cgt Arg	ggc Gly	gat Asp	tgg Trp 645	tgt Cys	gaa Glu	ctc Leu	gcc Ala	aag Lys 650	gag Glu	gtt Val	cag Gln	act Thr	aag Lys 655	gtt Val	1968		
gtc Val	gaa Glu	ata Ile	gta Val 660	ttg Leu	aag Lys	acg Thr	agt Ser	gag Glu 665	gtg Val	aac Asn	aag Lys	gct Ala	gta Val 670	gag Glu	tac Tyr	2016		
gtc Val	agg Arg	aag Lys 675	att Ile	gtg Val	aaa Lys	gag Glu	ttg Leu 680	gag Glu	gag Glu	ggc Gly	aag Lys	gtt Val 685	ccc Pro	ata Ile	gag Glu	2064		
Lys	Leu 690	Val	Ile	Trp	Lys	Thr 695	Leu	Ser	Lys	Arg	Leu 700	gag Glu	Glu	Tyr	Thr	2112		
Thr 705	Glu	Ala	Pro	His	Val 710	Val	Ala	Ala	Lys	Arg 715	Met	ctg Leu	Ser	Ala	Gly 720	2160		
Tyr	Arg	Val	Ser	Pro 725	Gly	Asp	Lys	Ile	Gly 730	Tyr	Val	ata Ile	Val	Lys 735	Gly	2208		
Gly	Gly	Arg	Ile 740	Ser	Gln	Arg	Ala	Trp 745	Pro	Tyr	Phe	atg Met	Val 750	Lys	Asp	2256		
Pro	Ser	Gln 755	Ile	Asp	Val	Thr	Tyr 760	Tyr	Val	Asp	His	caa Gln 765	Ile	Ile	Pro			
Ala	Ala 770	Leu	Arg	Ile	Leu	Gly 775	Tyr	Phe	Gly	Ile	Thr 780	gag Glu	Lys	Lys	Leu	2352		
aaa Lys 785	gca Ala	agt Ser	gca Ala	act Thr	999 Gly 790	cag Gln	aag Lys	act Thr	ctc Leu	ttc Phe 795	gac Asp	ttt Phe	cta Leu	gcc Ala	aag Lys 800	2400		

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4,7

aag agc aag taa Lys Ser Lys

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<212> PRT

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<400> 2

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Gly Lys Glu Pro Gln Val Ile Ile Trp Gly Ile Ala Glu Asn Gly Glu 20 25 30

Arg Val Val Leu Ile Asp Arg Ser Phe Arg Pro Tyr Phe Tyr Ala Leu 35 40 45

Leu Ala Pro Gly Ala Asp Pro Lys Gln Val Ala Gln Arg Ile Arg Ala 50 55 60

Leu Ser Arg Pro Lys Ser Pro Ile Ile Gly Val Glu Asp Asp Lys Arg 65 70 75 80

Lys Tyr Phe Gly Arg Pro Arg Arg Val Leu Arg Ile Arg Thr Val Leu 85 90 95

Pro Glu Ala Val Arg Glu Tyr Arg Glu Leu Val Lys Asn Val Asp Gly
100 105 110

Val Glu Asp Val Leu Glu Ala Asp Ile Arg Phe Ala Met Arg Tyr Leu 115 120 125

Ile Asp His Asp Leu Phe Pro Phe Thr Trp Tyr Arg Val Glu Ala Glu
130 135 140

Pro Leu Glu Asn Lys Met Gly Phe Arg Val Asp Lys Val Tyr Leu Val 145 150 155 160

Lys Ser Arg Pro Glu Pro Leu Tyr Gly Glu Ala Leu Ala Pro Thr Lys 165 170 175

Leu Pro Asp Leu Arg Ile Leu Ala Phe Asp Ile Glu Val Tyr Ser Lys

180 185 190

Gln Gly Ser Pro Arg Pro Glu Arg Asp Pro Val Ile Val Ile Ala Val Lys Thr Asp Asp Gly Asp Glu Val Leu Phe Ile Ala Glu Gly Lys Asp Asp Arg Lys Pro Ile Arg Glu Phe Val Glu Tyr Val Lys Arg Tyr Asp Pro Asp Ile Ile Val Gly Tyr Asn Asn Asn His Phe Asp Trp Pro Tyr 250 Leu Leu Arg Arg Ala Arg Ile Leu Gly Ile Lys Leu Asp Val Thr Arg 265 Arg Val Gly Ala Glu Pro Thr Thr Ser Val His Gly His Val Ser Val 280 Pro Gly Arg Leu Asn Val Asp Leu Tyr Asp Tyr Ala Glu Glu Met Pro 295 300 Glu Ile Lys Ile Lys Ser Leu Glu Glu Val Ala Glu Tyr Leu Gly Val 310 Met Lys Lys Ser Glu Arg Val Ile Ile Asn Trp Trp Glu Ile Pro Asp 325 Tyr Trp Asp Asp Pro Lys Lys Arg Pro Leu Leu Gln Tyr Ala Arg 345 Asp Asp Val Arg Ala Thr Tyr Gly Leu Ala Glu Lys Ile Leu Pro Phe 360 365 Ala Ile Gln Leu Ser Tyr Val Thr Gly Leu Pro Leu Asp Gln Val Gly 370 375 380 Ala Met Ser Val Gly Phe Arg Leu Glu Trp Tyr Leu Ile Arg Ala Ala 385 390 395 Phe Lys Met Lys Glu Leu Val Pro Asn Arg Val Glu Arg Pro Glu Glu 405 410 415

Thr Tyr Arg Gly Ala Ile Val Leu Glu Pro Leu Arg Gly Val His Glu
420 425 430

Asn Ile Ala Val Leu Asp Phe Ser Ser Met Tyr Pro Asn Ile Met Ile 435 440 445

Lys Tyr Asn Val Gly Pro Asp Thr Leu Val Arg Pro Gly Glu Lys Cys 450 455 460

Gly Glu Cys Gly Cys Trp Glu Ala Pro Glu Val Lys His Arg Phe Arg 465 470 475 480

Arg Cys Pro Pro Gly Phe Phe Lys Thr Val Leu Glu Arg Leu Leu Glu 485 490 495

Leu Arg Lys Arg Val Arg Ala Glu Met Lys Lys Tyr Pro Pro Asp Ser 500 510

Pro Glu Tyr Arg Leu Leu Asp Glu Arg Gln Lys Ala Leu Lys Val Leu 515 520 525

Ala Asn Ala Ser Tyr Gly Tyr Met Gly Trp Ser Gly Ala Arg Trp Tyr 530 540

Cys Arg Glu Cys Ala Lys Ala Val Thr Ala Trp Gly Arg His Leu Ile 545 550 555 560

Arg Thr Ala Ile Asn Ile Ala Arg Lys Leu Gly Leu Lys Val Ile Tyr 565 570 575

Gly Asp Thr Asp Ser Leu Phe Val Thr Tyr Asp Pro Glu Lys Val Glu 580 585 590

Asn Phe Ile Lys Ile Ile Lys Glu Glu Leu Gly Phe Glu Ile Lys Leu 595 600 605

Glu Lys Val Tyr Lys Arg Leu Phe Phe Thr Glu Ala Lys Lys Arg Tyr 610 615 620

Ala Gly Leu Leu Glu Asp Gly Arg Ile Asp Ile Val Gly Phe Glu Ala 625 630 635 640 Val Arg Gly Asp Trp Cys Glu Leu Ala Lys Glu Val Gln Thr Lys Val 645 650 655

Val Glu Ile Val Leu Lys Thr Ser Glu Val Asn Lys Ala Val Glu Tyr 660 665 670

Val Arg Lys Ile Val Lys Glu Leu Glu Glu Gly Lys Val Pro Ile Glu 675 680 685

Lys Leu Val Ile Trp Lys Thr Leu Ser Lys Arg Leu Glu Glu Tyr Thr 690 695 700

Thr Glu Ala Pro His Val Val Ala Ala Lys Arg Met Leu Ser Ala Gly 705 710 715 720

Tyr Arg Val Ser Pro Gly Asp Lys Ile Gly Tyr Val Ile Val Lys Gly 725 730 735

Gly Gly Arg Ile Ser Gln Arg Ala Trp Pro Tyr Phe Met Val Lys Asp 740 745 750

Pro Ser Gln Ile Asp Val Thr Tyr Tyr Val Asp His Gln Ile Ile Pro 755 760 765

Ala Ala Leu Arg Ile Leu Gly Tyr Phe Gly Ile Thr Glu Lys Lys Leu 770 775 780

Lys Ala Ser Ala Thr Gly Gln Lys Thr Leu Phe Asp Phe Leu Ala Lys 785 790 795 800

Lys Ser Lys